

# **Product datasheet for TP710215**

### OriGene Technologies, Inc.

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## IDH2 (NM\_002168) Human Recombinant Protein

#### **Product data:**

**Product Type:** Recombinant Proteins

**Description:** Purified recombinant protein of mutant(R172M) of Homo sapiens isocitrate dehydrogenase 2

(NADP+), mitochondrial (IDH2), with C-terminal DDK tag, expressed in sf9 cells, 20 µg

Species: Human

**Expression Host:** Sf9

**Expression cDNA Clone** 

or AA Sequence:

A DNA sequence from TrueORF clone, RC400102, encoding human full-length IDH2 R172M

Tag: C-DDK

**Predicted MW:** 46.6 kDa

**Concentration:** >0.05 μg/μL as determined by microplate BCA method

**Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining

**Buffer:** 50 mM Tris-HCl, 100 mM glycine, pH 8.0, 10% glycerol

**Note:** For testing in cell culture applications, please filter before use. Note that you may experience

some loss of protein during the filtration process.

Storage: Store at -80°C.

Stability: Stable for 12 months from the date of receipt of the product under proper storage and

handling conditions. Avoid repeated freeze-thaw cycles.

RefSeq: NP 002159

 Locus ID:
 3418

 UniProt ID:
 P48735

 RefSeq Size:
 1740

 Cytogenetics:
 15q26.1

RefSeq ORF: 1356

Synonyms: D2HGA2; ICD-M; IDH; IDHM; IDP; IDPM; mNADP-IDH



**Summary:** 

Isocitrate dehydrogenases catalyze the oxidative decarboxylation of isocitrate to 2-oxoglutarate. These enzymes belong to two distinct subclasses, one of which utilizes NAD(+) as the electron acceptor and the other NADP(+). Five isocitrate dehydrogenases have been reported: three NAD(+)-dependent isocitrate dehydrogenases, which localize to the mitochondrial matrix, and two NADP(+)-dependent isocitrate dehydrogenases, one of which is mitochondrial and the other predominantly cytosolic. Each NADP(+)-dependent isozyme is a homodimer. The protein encoded by this gene is the NADP(+)-dependent isocitrate dehydrogenase found in the mitochondria. It plays a role in intermediary metabolism and energy production. This protein may tightly associate or interact with the pyruvate dehydrogenase complex. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Feb 2014]

**Protein Pathways:** 

Citrate cycle (TCA cycle), Glutathione metabolism, Metabolic pathways

# **Product images:**

